



# TIN-X v2: modernized architecture with REST API for sustainability & interoperability

Jeremy Yang<sup>1</sup>, Daniel Cannon<sup>2</sup>, Cristian Bologa<sup>1</sup>, Stephen Mathias<sup>1</sup>, Lars Juhl Jensen<sup>3</sup>, Stephan Schürer<sup>4</sup>, Dušica Vidović<sup>4</sup>, and Tudor Oprea<sup>1</sup> <sup>1</sup>University of New Mexico, Albuquerque, NM, USA; <sup>2</sup>Iterative Consulting, LLC, Albuquerque, NM, USA; <sup>3</sup>Novo Nordisk Center for Protein Research, University of Denmark, Copenhagen, Denmark; <sup>4</sup>University of Miami, Miami, FL, USA

### **Overview**

- TIN-X = Target Importance and Novelty Explorer
- Bibliometric analytical method for prioritizing targets
- Data source: TCRD/Pharos
- **Prototype** developed into **product -- DevOps**
- Proposed informatics tools exemplar

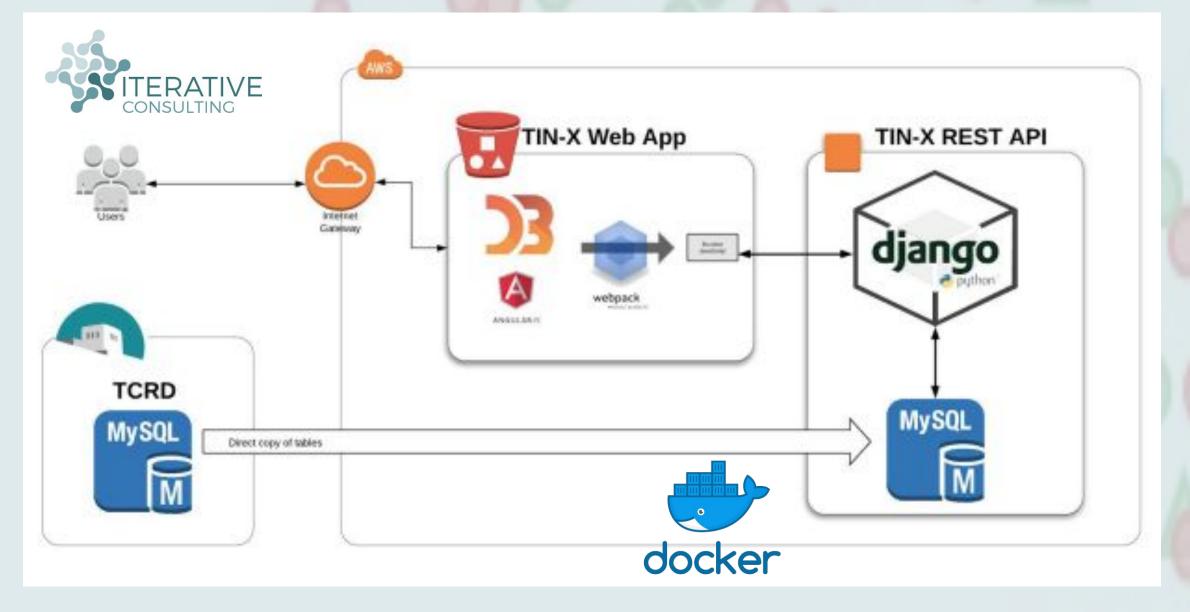


# **NEW: REST API, Swagger powered, public**

N-X REST API	TIN-2	X REST API	<pre># Install the \$ pip install</pre>	TIN-X REST API
rticles	articles         list         Image: Construction of the database         Image: Construction of the database         Description         Limit       Number of results to return per page.         Image: Construction of the database         Image: Construction of the database		<pre>\$ pip instatt # Load the sch \$ coreapi get # Interact wit \$ coreapi acti</pre>	Api Root Api Root Api Root This is the root of the TIN-X REST API. Explore the API by following the links below. For more detailed information, please consult the API Documentation GET / HTTP 200 OK Allow: GET, HEAD, OPTIONS Content-Type: application/json Vary: Accept { "diseases": "http://api.newdrugtargets.org/diseases/", "targets": "http://api.newdrugtargets.org/targets/", "articles": "http://api.newdrugtargets.org/articles/", "dto": "http://api.newdrugtargets.org/dto/"
	title journal author			api.newdrugtargets.org

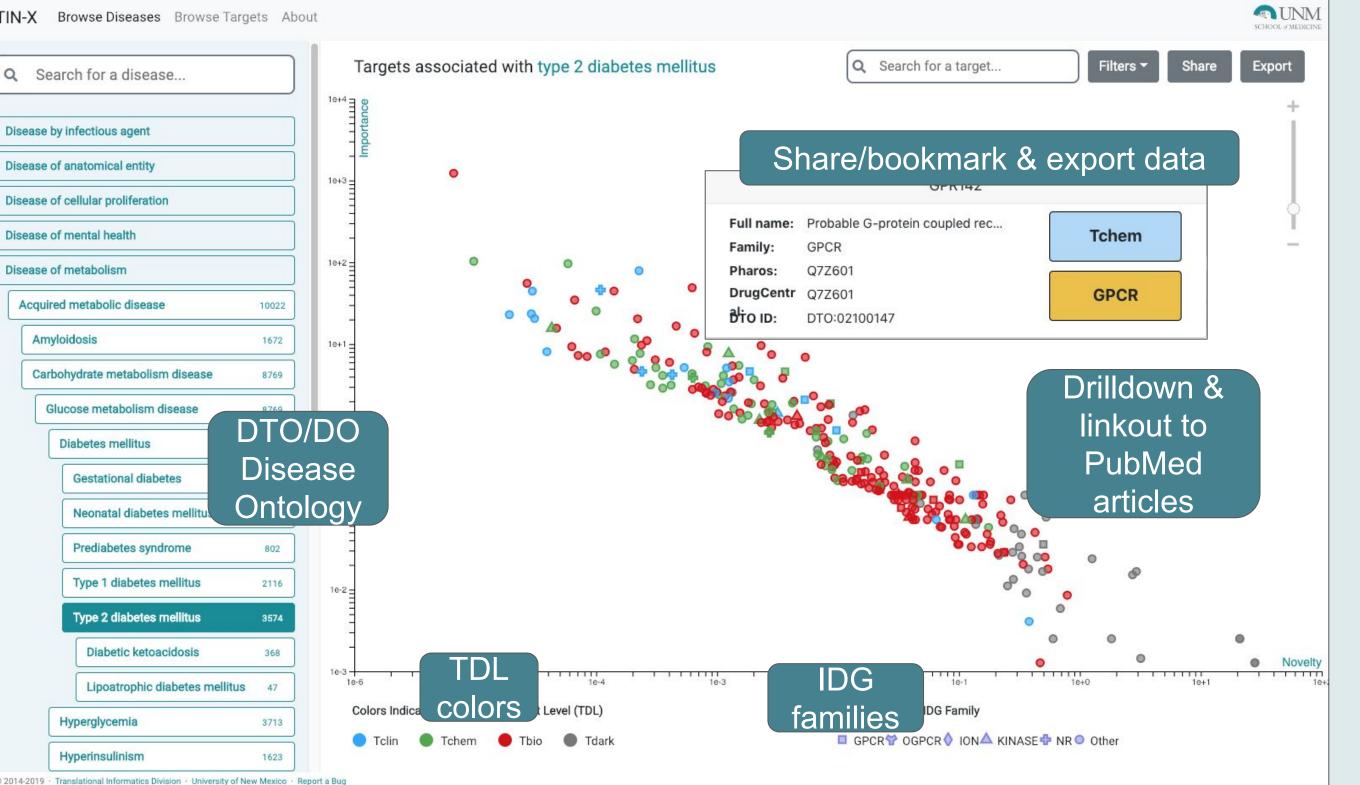
- New architecture for sustainability
- New REST API for *interoperability*
- Version 2.0 released Feb 18, 2019

# **New Architecture, TIN-X Version 2.0 (newdrugtargets.org)**



- Thin, modern, responsive web app
- Tighter TCRD integration via MySQL
- Better automated for faster PubMed updates
- Dockerized for rapid, flexible, economical deployment
- Design patterned for replication and reuse.

## TIN-X v2 for Type 2 diabetes mellitus

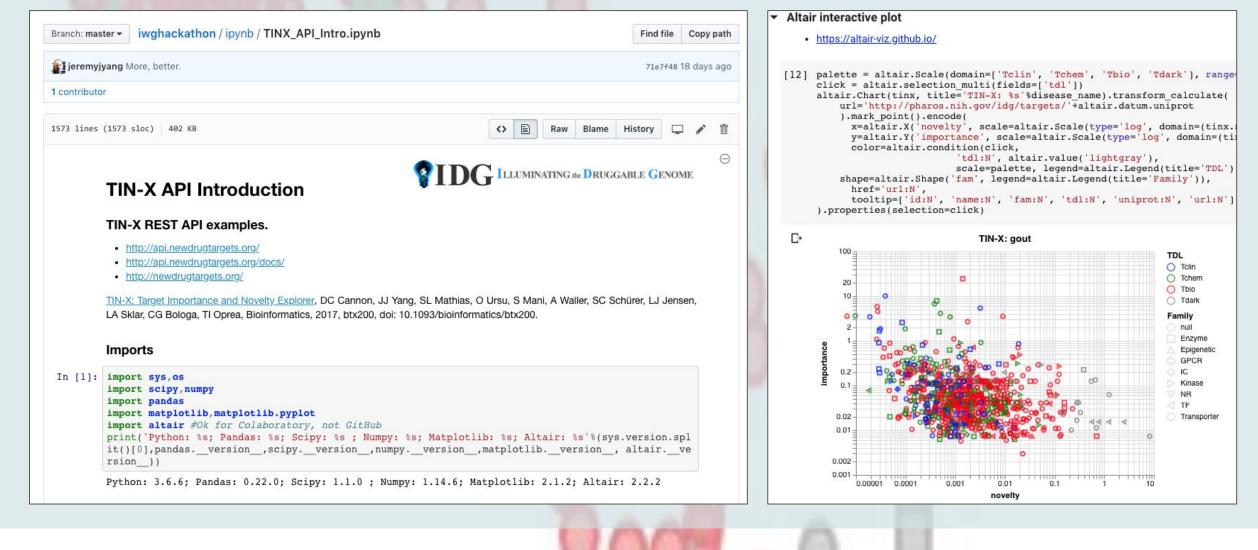


# History, Background, Contributions

TIN-X is designed to prioritize and visualize associations between proteins and diseases, from scientific literature (PubMed) text mining by JensenLab, via TCRD, and organized by Drug Target Ontology (DTO) based disease and protein classifications. TIN-X was initially conceived and prototyped by Cristian Bologa, then engineered as a full stack webapp by Daniel Cannon, deployed via AWS. Motivated by its success and perceived value to researchers, TIN-X has been continually maintained, updated, and improved. Recently, TIN-X has undergone a major revision to version 2.0, designed and implemented by Iterative Consulting, LLC, co-founded by Daniel Cannon. The new architecture conforms to modern software engineering standards, includes a Swagger/Django REST API, D3 thin client, and tight integration with TCRD. Updates and deployment automation employs Docker and AWS (EC2, S3, CloudFront). Source code is managed via Bitbucket and GitHub. The improvements address the Resource Sharing Plan of KMC, and NIH policies and principles concerning digital resource sharing (e.g. FAIR) as emphasized by the NIH Strategic Plan for Data Science. Specifically, the new architecture

- Facilitates timely updates.
- Facilitates code support, maintenance, and further development.
- Facilitates clients with API access to TIN-X results.
- Provides transparency and reproducibility of workflow protocols.
- Serves as exemplar for IDG KMC based informatics and analytics tools. In addition, the development and evolution of TIN-X from research prototype to enterprise component is a proposed DevOps design pattern.





#### References:

- 1. TIN-X: Target Importance and Novelty Explorer, DC Cannon, JJ Yang, SL Mathias, O Ursu, S Mani, A Waller, SC Schürer, LJ Jensen, LA Sklar, CG Bologa, TI Oprea, Bioinformatics, 2017, btx200, doi: 10.1093/bioinformatics/btx200.
- 2. NIH Strategic Plan for Data Science, https://grants.nih.gov/grants/rfi/NIH-Strategic-Plan-for-Data-Science.pdf.

This work was supported by US National Institutes of Health (grants U54 CA189205 and U24 224370), Illuminating the Druggable Genome Knowledge Management Center (IDG KMC),

